AN   1996:491081 CAPLUS
DN   125:161780
TI   Cloning of a novel ubiquitin-conjugating enzyme (E2) gene from the ciliate
     Paramecium tetraurelia
AU   Okano, Satoshi; Tokushima, Hideyuki; Nakaoka, Yasuo; Shimizu, Kikuo
CS   Department of Biophysical Engineering, Faculty of Engineering Science,
     Osaka University, 1-3 Machikaneyama, Toyonaka, Osaka, 560, Japan
SO   FEBS Letters (1996), 391(1,2), 1-4
     CODEN: FEBLAL; ISSN: 0014-5793
PB   Elsevier
DT   Journal
LA   English
AB   We isolated a 1.7 kb gene (UbcP1) for a ubiquitin-conjugating enzyme from
     a P. tetraurelia cDNA library and sequenced it. Its deduced polypeptide
     sequence consists of 425 amino acid residues (48 kDa). The UbcP1 protein
     contains novel N- and C-terminal extensions in addition to a UBC domain, and
     within the UBC domain it shares low identity with sequences of other known
     E2s. A constructed phylogenetic tree suggests that the UbcP1 protein may
     represent a member of a distinct subfamily of E2s. Southern blot anal.
     showed that the N-terminal extension of the UbcP1 is conserved in P.
     multimicronucleatum.
IT   180391-20-2
     RL: BOC (Biological occurrence); BSU (Biological study, unclassified); PRP
     (Properties); BIOL (Biological study); OCCU (Occurrence)
        (amino acid sequence; cloning of novel ubiquitin-conjugating enzyme
        (E2) gene from ciliate Paramecium tetraurelia)
RN   180391-20-2 CAPLUS
CN   Enzyme, E2 (ubiquitin-carrier) (Paramecium tetraurelia strain 51s gene
     UbcP1) (9CI) (CA INDEX NAME)

SEQ   1 MGGMYFKEEE LSRIGAVNAF FSAFADKTLA FEFNHIVKLV WFESFITDKC
        51 DFTNDFNNFI KLVDDASPRG GTKCYDAIAY AIEQLKEIKK KYPNIILRII
       101 ALTDGDDNQS KENPQSLVNR IFENQIIIDS FVVNNDCVGL KTLTHATNGR
       151 CYCPQTLAEG MSLFEIESIL SISHREQKEY PKELQDLNSL KDKPFDTDGM
       201 KVVSMDVQKI AVMKKEEILK KISQIESAPQ NSSSTSQSVN NTARILKELQ
       251 DVTAQGDKLN FKCYPTADDI KTWKILLYGP KGTVYEGGLY ILSYVFTQNY
       301 PFRPPKVQFI TKLYHPNVSR GGSLCLDVLN TSWSPLLTTT KVLDAVSVML
       351 QNPNADDALD CNIAAIYKHE PELFKQNALK EKLEAASPSE DNLLADILGA
       401 VDINSQEYLD TKKELQTWLE YQKSN

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     FILE 'CAPLUS' ENTERED AT 12:47:07 ON 01 NOV 2005
            S QEHIGV/SQSP OR EHIGVM/SQSP OR HIGVMK/SQSP OR IGVMKK/SQSP OR G

     FILE 'REGISTRY' ENTERED AT 12:48:30 ON 01 NOV 2005
L1   60 S VMKKEE/SQSP

     FILE 'CAPLUS' ENTERED AT 12:48:56 ON 01 NOV 2005
L2   32 S L1

     FILE 'REGISTRY' ENTERED AT 12:48:58 ON 01 NOV 2005
L3   37 S GVMKKE/SQSP

     FILE 'CAPLUS' ENTERED AT 12:49:00 ON 01 NOV 2005
L4   26 S L3

L14　　11 SEA L13 AND PY<2001
　　　　D 1-11 TI
　　　　D 1-11 BIB AB HITSEQ

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             FILE 'REGISTRY' ENTERED AT 12:49:02 ON 01 NOV 2005
L5              36 S IGVMKK/SQSP

             FILE 'CAPLUS' ENTERED AT 12:49:04 ON 01 NOV 2005
L6              22 S L5

             FILE 'REGISTRY' ENTERED AT 12:49:05 ON 01 NOV 2005
L7              22 S HIGVMK/SQSP

             FILE 'CAPLUS' ENTERED AT 12:49:08 ON 01 NOV 2005
L8              12 S L7

             FILE 'REGISTRY' ENTERED AT 12:49:08 ON 01 NOV 2005
L9              25 S EHIGVM/SQSP

             FILE 'CAPLUS' ENTERED AT 12:49:11 ON 01 NOV 2005
L10             15 S L9

             FILE 'REGISTRY' ENTERED AT 12:49:12 ON 01 NOV 2005
L11             22 S QEHIGV/SQSP

             FILE 'CAPLUS' ENTERED AT 12:49:14 ON 01 NOV 2005
L12             12 S L11
L13             60 S L12 OR L10 OR L8 OR L6 OR L4 OR L2
L14             11 S L13 AND PY<2001

=> d his full

        (FILE 'HOME' ENTERED AT 12:47:02 ON 01 NOV 2005)

        FILE 'CAPLUS' ENTERED AT 12:47:07 ON 01 NOV 2005
                S QEHIGV/SQSP OR EHIGVM/SQSP OR HIGVMK/SQSP OR IGVMKK/SQSP OR G

        FILE 'REGISTRY' ENTERED AT 12:48:30 ON 01 NOV 2005
L1           60 SEA VMKKEE/SQSP

        FILE 'CAPLUS' ENTERED AT 12:48:56 ON 01 NOV 2005
L2           32 SEA L1

        FILE 'REGISTRY' ENTERED AT 12:48:58 ON 01 NOV 2005
L3           37 SEA GVMKKE/SQSP

        FILE 'CAPLUS' ENTERED AT 12:49:00 ON 01 NOV 2005
L4           26 SEA L3

        FILE 'REGISTRY' ENTERED AT 12:49:02 ON 01 NOV 2005
L5           36 SEA IGVMKK/SQSP

        FILE 'CAPLUS' ENTERED AT 12:49:04 ON 01 NOV 2005
L6           22 SEA L5

        FILE 'REGISTRY' ENTERED AT 12:49:05 ON 01 NOV 2005
L7           22 SEA HIGVMK/SQSP

        FILE 'CAPLUS' ENTERED AT 12:49:08 ON 01 NOV 2005
L8           12 SEA L7

        FILE 'REGISTRY' ENTERED AT 12:49:08 ON 01 NOV 2005
L9           25 SEA EHIGVM/SQSP

        FILE 'CAPLUS' ENTERED AT 12:49:11 ON 01 NOV 2005
L10          15 SEA L9

        FILE 'REGISTRY' ENTERED AT 12:49:12 ON 01 NOV 2005
L11          22 SEA QEHIGV/SQSP

        FILE 'CAPLUS' ENTERED AT 12:49:14 ON 01 NOV 2005
L12          12 SEA L11
L13          60 SEA L12 OR L10 OR L8 OR L6 OR L4 OR L2
```

L14 ANSWER 11 OF 11 CAPLUS COPYRIGHT 2005 ACS on STN
AN   1996:491081 CAPLUS
DN   125:161780
TI   Cloning of a novel ubiquitin-conjugating enzyme (E2) gene from the ciliate
     Paramecium tetraurelia
AU   Okano, Satoshi; Tokushima, Hideyuki; Nakaoka, Yasuo; Shimizu, Kikuo
CS   Department of Biophysical Engineering, Faculty of Engineering Science,
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SO   FEBS Letters (1996), 391(1,2), 1-4
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     within the UBC domain it shares low identity with sequences of other known
     E2s. A constructed phylogenetic tree suggests that the UbcP1 protein may
     represent a member of a distinct subfamily of E2s. Southern blot anal.
     showed that the N-terminal extension of the UbcP1 is conserved in P.
     multimicronucleatum.
IT   180391-20-2
     RL: BOC (Biological occurrence); BSU (Biological study, unclassified); PRP
     (Properties); BIOL (Biological study); OCCU (Occurrence)
         (amino acid sequence; cloning of novel ubiquitin-conjugating enzyme
         (E2) gene from ciliate Paramecium tetraurelia)
RN   180391-20-2 CAPLUS
CN   Enzyme, E2 (ubiquitin-carrier) (Paramecium tetraurelia strain 51s gene
     UbcP1) (9CI)   (CA INDEX NAME)

SEQ   1 MGGMYFKEEE LSRIGAVNAF FSAFADKTLA FEFNHIVKLV WFESFITDKC
        51 DFTNDFNNFI KLVDDASPRG GTKCYDAIAY AIEQLKEIKK KYPNIILRII
       101 ALTDGDDNQS KENPQSLVNR IFENQIIIDS FVVNNDCVGL KTLTHATNGR
       151 CYCPQTLAEG MSLFEIESIL SISHREQKEY PKELQDLNSL KDKPFDTDGM
       201 KVVSMDVQKI AVMKKEEILK KISQIESAPQ NSSSTSQSVN NTARILKELQ
       251 DVTAQGDKLN FKCYPTADDI KTWKILLYGP KGTVYEGGLY ILSYVFTQNY
       301 PFRPPKVQFI TKLYHPNVSR GGSLCLDVLN TSWSPLLTTT KVLDAVSVML
       351 QNPNADDALD CNIAAIYKHE PELFKQNALK EKLEAASPSE DNLLADILGA
       401 VDINSQEYLD TKKELQTWLE YQKSN

=> d his

     (FILE 'HOME' ENTERED AT 12:47:02 ON 01 NOV 2005)

     FILE 'CAPLUS' ENTERED AT 12:47:07 ON 01 NOV 2005
               S QEHIGV/SQSP OR EHIGVM/SQSP OR HIGVMK/SQSP OR IGVMKK/SQSP OR G

     FILE 'REGISTRY' ENTERED AT 12:48:30 ON 01 NOV 2005
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L2   32 S L1

     FILE 'REGISTRY' ENTERED AT 12:48:58 ON 01 NOV 2005
L3   37 S GVMKKE/SQSP

     FILE 'CAPLUS' ENTERED AT 12:49:00 ON 01 NOV 2005
L4   26 S L3

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 201 DIDPLTTLKY ILSLNDQLYD TADVILLEVI HLFKKLYIPH VFDDSYLLIN
 251 DEDKDDDDEN NDENNDKESD EEDDHGDDHN DYNINSNLKE EEKYKYNYND
 301 PIYDENKNNK ENNLLDYKKI NNIHNITSNE RYSQSTNHIY TQNNKIGDTY
 351 KYISNKSNNV IHHKNEYKEL GEELNLLSED INFKQKKLNE NNYNQYKNHV
 401 IKILLNMLSK NVSYSVLYEG ACCLLYMSTS ALSIKTACEC FIKLLINQHD
 451 NNIKLIVIDK LYYIMCKWKN ILENYAMDLL RALNFPSRDI KVKILNLVLH
 501 VLTKRNVHLV LNVLKKELLK LNDQVVYTKH IITTTNNNNN NTNMNMSNNN
 551 NDNNDNNNNN NNNNNNNNNN NNSNNNSSNN NNNNLSTNYQ EVTSYKKILI
 601 KSLQHICNLY SSECLHIVDL LLIYVNHNEQ DINYEAAVCI RKLVKNNNFQ
 651 NNILEKMIDS IFDIKKATIL RIFFWVIGQY MFNEHMIINF LNNLYDNLSP
 701 LLNNSLENDM INKIQNERFK KNGNINFNLS NSNIQTKTVI LEDGTYATEA
 751 FLKNQNICNK NNDKNKENTF VYNILYENDD ILLSVICVSL TKLYLKLLSK
 801 MNNSFDFILN QNEIICSKEK KEMKHSLIKH EHFLNHVTNN NTLDTEECVI
 851 SKDPNRISTN SENKITEHDI NEYRNKCIYI LASIIKYIGE KNSKPDSNNI
 901 YEYDSNVIRI NQCLKIFFYL TANKKELDEQ SKKLIKIFID GDYYYNKFLE
 951 KEEEKYCSIY GSNYKYMKRN NMYANKTNGG NKTFDGNNTF DGNNTFDGNN
1001 TFDGNNILEK STYSRNKLFL NDQGEKENVD DDIYFRVLKE KKNILNMLDE
1051 ETSSVCINEN LEKLKLKYTL NNDMLFEENE FQYPSIKYNY SSLFLAKLYN
1101 SQILTGTDDD IFIEALPIIS NVNLIIEIYV YNQSNVYLQN IYINLSTHGN
1151 LKPIDKIPEF NLAPNEKKKF KISVKVHTTE AGIIFGYVFY ERKNDNHKNY
1201 IVLNELHINM TDYINASFIS SHLFRIMWSE FEWENKINIH TSIRDAFELL
1251 KLIIKNTNMT IVERFMPLEY YDMEIKNNVN QVNISPIDIY ISYISTLEDL
1301 KCLVNNSAFF SVNLFSRSIF GEDSLANLSV QKSADGKLSG SIRVRSRTQV
1351 RKNKTKQNKT KQNKPDGYIT LLYGIALSLG DKLTLLQTGI NADM
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`.CNT   30   THERE ARE 30 CITED REFERENCES AVAILABLE FOR THIS RECORD
                ALL CITATIONS AVAILABLE IN THE RE FORMAT

D L2 20 TI BIB HITSEQ

ANSWER 20 OF 21   CAPLUS   COPYRIGHT 2005 ACS on STN
Localization and function of five glutamate transporters cloned from the
salamander retina
1998:369436   CAPLUS
129:146956
Localization and function of five glutamate transporters cloned from the
salamander retina

Eliasof, Scott; Arriza, Jeffrey L.; Leighton, Barbara H.; Amara, Susan G.;
Kavanaugh, Michael P.
Vollum Institute L-474, Oregon Health Sciences University, Portland, OR,
97210, USA

203212-47-9, Amino acid transporter, glutamate-transporting 2B
(Ambystoma tigrinum gene sEAAT2B)
RL: BOC (Biological occurrence); BPR (Biological process); BSU (Biological
study, unclassified); PRP (Properties); BIOL (Biological study); OCCU
(Occurrence); PROC (Process)
     (amino acid sequence; localization and function of five glutamate
     transporters cloned from the salamander retina)
203212-47-9 CAPLUS
Glutamate transporter 2B (Ambystoma tigrinum gene sEAAT2B) (9CI)   (CA
INDEX NAME)

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  1 MTNNVEIPLS DGHHHHLALE SLPIKDCCSK RCGWLTRNLL LILTIGGVIL
 51 GSVFGTLLRL LPPLDSNIVL LISFPGDILM RMLKMLILPL VISSLISGLA
101 GLDAKSSGRM GTRAMVYYMS TTILAAVLGV ILVLSIHPGN PKLKKQVSIS
151 TKNEEVSSLD AFLDLIRNLF PENLVQACFQ QIQTVSKKVP VTPSPHELLN
201 QVNISVSQSN SSMLNSTTAE GNAGPTMVTQ KKLEFKAGMN VLGLIGFFIA
251 FGIAMGKMGE QARLMADFFN ILNEIIMKLV NMIMWYSPFG IACLICGKIA
301 AIKDLEVVAR QLGMYMVTVI VGLVIHGGIV LPLIFFSITR TNPYTFYGGI
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351 FQAWITALGT ASSAGTLPVT FRCLEENLKI ·DKRVTRFVLP IGATINMDGT
401 ALYEAVAAIF IAQMNDVSLD GGQIATVSLT ATLASVGAAS IPSAGLVTML
451 LILTAVGLPT QDISLLIAVD WLLDRMRTSI NVVGDSFGAG IIYHLSKAEL
501 AAIDAKHEAS KEGGKLHLMN HEAAQKHPSD GMSSPPFSGY TSLPTYEYEA
551 HEKQESLFGH EDDGEEPWST SRNSNEKKME D
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RE.CNT   35   THERE ARE 35 CITED REFERENCES AVAILABLE FOR THIS RECORD
               ALL CITATIONS AVAILABLE IN THE RE FORMAT